

Amendment

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Serial No.: 10/516,578

Confirmation No.: 5513

Filed: November 16, 2005

For: PSEUDOTYPED RETROVIRUSES

Amendments to the Specification

Please amend the paragraph beginning at page 6, line 17, as follows:

Filoviral proteins, such as those of Ebola virus (e.g., Ebola Zaire [complete genome, GenBank Acc. No. NC 002549; glycoprotein, ~~GenBank~~ GenBank Acc. No. U23187], Ebola Reston [complete genome, GenBank Acc. No. NC 004161; glycoprotein, GenBank Acc. No. U23152] and Ebola Sudan [glycoprotein, GenBank Acc. No. U23069]; see Fig. 1) and Marburg virus [complete genome, GenBank Acc. No. NC 001608; glycoprotein, GenBank Acc. No. Z12132]), have *O*-glycosylation regions and infect a broad spectrum of mammalian hosts. By way of non-limiting example, the *O*-glycosylation region for the Ebola glycoprotein (Zaire) is from about ~~nucleotide-codon~~ 309 to ~~nucleotide-codon~~ 489 of SEQ ID NO:1. The nucleotide sequences encoding the filoviral glycoproteins may be obtained as described in Sanchez et al. (1993) *Virus Res.* 29 (3):215-240 and Will et al., (1993) *J. Virol.* 67:1203-1210. It is expected that other viruses not specifically mentioned herein and having glycoproteins of similar structure to the filoviral glycoproteins may be advantageously used in the present invention.